Serial Number: 09/101,423B

CRF Processing Date: 1/11/2001
Edited by: _____
Verified by: _____ (STIC staff)

ENTERED

RECEIVED
JAN 26 2004
TECH CENTER 1600/2900

- ☐ Changed a file from non-ASCII to ASCII

- ☐ Changed the margins in cases where the sequence text was wrapped down to the next line.

- ☐ Edited a format error in the Current Application Data section, specifically: _____

- ☑ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____

- ☐ Added the mandatory heading and subheadings for "Current Application Data".

- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

- ☐ Inserted colons after headings/subheadings. Headings edited included: _____

- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

- ☑ Deleted: ☑ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____

- ☐ Inserted mandatory headings, specifically: _____

- ☐ Corrected an obvious error in the response, specifically: _____

- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

- ☐ Corrected an error in the Number of Sequences field, specifically: _____

- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____

- ☐ Other: _____

Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/101,423B

DATE: 01/11/2001
TIME: 13:19:54

Input Set : A:\Sequence Listing
Output Set: N:\CRF3\01112001\I101423B.raw

Does Not Comply
Corrected Diskette Needed

```
       3 <110> APPLICANT: Rudland, Philip S.
       4        Barraclough, Roger B.
W--> 5 <120> TITLE OF INVENTION: Metastasis Inducing DNA's
W--> 6 <130> FILE REFERENCE: WPT 0114 PUS
W--> 7 <140> CURRENT APPLICATION NUMBER: US 09/101,423B
       8 <141> CURRENT FILING DATE: 1998-11-27
       9 <150> PRIOR APPLICATION NUMBER: PCT/GB97/00074
      10 <151> PRIOR FILING DATE: 1997-01-10
W--> 11 <160> NUMBER OF SEQ ID: 8


ERRORED SEQUENCES

     271 <210> SEQ ID NO: 8
     272 <211> LENGTH: 47 base pairs
     273 <212> TYPE: DNA
     274 <213> ORGANISM: Homo sapiens
W--> 275 <400> SEQUENCE: 8
     277 agctataatg cggccgcata ttcggcctga tcggccgcaa gcttgga     47
W--> 279 M:\clients\W\WPT\0114pusa\2nd revis seq list 12-20-00.wpd
E--> 280 kathy 12/20/00
```

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/101,423B

DATE: 01/11/2001
TIME: 13:19:56

Input Set : A:\Sequence Listing
Output Set: N:\CRF3\01112001\I101423B.raw

```
L:5   M:283 W: Missing Blank Line separator, <120> field identifier
L:6   M:283 W: Missing Blank Line separator, <130> field identifier
L:7   M:283 W: Missing Blank Line separator, <140> field identifier
L:11  M:283 W: Missing Blank Line separator, <160> field identifier
L:59  M:283 W: Missing Blank Line separator, <400> field identifier
L:275 M:283 W: Missing Blank Line separator, <400> field identifier
L:279 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:5
L:280 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:8
L:280 M:254 E: No. of Bases conflict, LENGTH:Input:0 Counted:54 SEQ:8
L:280 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:6
L:280 M:112 C: (48) String data converted to lower case,
L:280 M:252 E: No. of Seq. differs, <211>LENGTH:Input:47 Found:54 SEQ:8
```